

[illegible]

em.pat:EO9533	+ 1197.00	1983.25	1.6e-102	2055	EO9533 CDNA encoding human
gb.pat:A36817	+ 1197.00	1983.17	1.6e-102	2070	A36817 Sequence 13 from Pat
gb.ro:NM05BMPRC	+ 1197.00	1982.07	1.8e-102	2292	D168250 mouse mRNA for BMP 1
gb.ro:NM046672	+ 1197.00	1981.56	2.0e-102	2402	D046672 Mus musculus type I
gb.ro:RATPBA41	+ 1194.00	1980.94	2.1e-102	1599	D17667 Rat mRNA for bone me
gb.prl:H0MSKRN	+ 1192.00	1980.74	4.7e-102	2085	D02911 Human novel serie k
gb.ro:RATRALX3	+ 1194.00	1974.10	5.1e-102	3003	D33082 Rat mRNA for bone m
gb.ro:R75359	+ 1189.00	1973.53	6.8e-102	1699	D75359 bone morphogenetic p
gb.com:RTU5B095	+ 1194.00	1971.89	6.8e-102	2715	U5B095 Bos taurus actin r
gb.prl:H5ALKX2A	+ 1192.00	1971.81	6.8e-102	2724	Z22534 H.sapiens ALK-2 mRNA
gb.pat:A33807	+ 1181.50	1955.14	5.8e-101	2494	A38807 Sequence 3 from Pat
gb.ov:AB011826	+ 1176.00	1949.58	1.2e-100	1780	AB011826 Danio rerio BMP-1
gb.ro:RATACTIEEC	+ 1175.00	1949.54	1.2e-100	1530	D123448 Sequence 1 from pate
gb.pat:125007	+ 1175.00	1949.54	1.2e-100	1530	D125007 Sequence 1 from pate
gb.ro:NM05SK7L	+ 1172.00	1941.30	3.4e-100	2056	D04672 Mus musculus N1H3T3
gb.ov:MG004673	+ 1166.50	1933.20	3.5e-99	3251	U04672 Gallus gallus type I
gb.ov:GV036622	+ 1166.50	1933.54	3.5e-99	3405	U03622 Gallus laevis activ
gb.ov:XU045914	+ 1160.50	1919.50	7.5e-98	3079	AB001073 Xenopus laevis mRN
gb.ov:AF0001073	+ 1160.50	1899.53	5.9e-97	3747	AF038425 Danio rerio activ1
gb.ov:AF0384425	+ 1140.00	1883.23	5.9e-97	3079	AF038425 Xenopus laevis ALK
gb.ro:NM05ALK1B	+ 1134.50	1878.89	1.0e-96	1962	D48015 Mus musculus activ
gb.ro:NM05ALK1B	+ 1126.50	1865.76	5.5e-96	1909	D38164 M.musculus (NH Swis
gb.pat:A38815	+ 1122.50	1851.25	3.6e-95	1922	A38815 Sequence 11 from Pat
gb.pat:123850	+ 1114.00	1847.27	5.9e-95	1518	D136088 Rattus norvegicus (C
gb.prl:123509	+ 1114.00	1847.27	5.9e-95	1518	D123850 Sequence 3 from pate
gb.prl:HSALK1A	+ 1091.50	1806.65	1.1e-92	1970	Z22533 H.sapiens ALK-1 mRNA
gb.pat:AA806074	+ 1121.50	1806.65	1.1e-92	204517	AC006074 Drosophila melan
gb.pat:A38805	+ 1091.50	1806.62	1.1e-92	1984	A38805 Sequence 1 from Pate
gb.prl:HM05FBRS	+ 1090.50	1806.41	1.1e-92	1733	L17075 Human TGF-b superfam
gb.inl:DM008K2A	+ 1090.50	1800.42	2.4e-92	3009	D33764 Drosophila melanogas
gb.inl:DM011442	+ 1090.50	1799.23	3.2e-92	3682	D11442 Drosophila melanogas
gb.inl:DM08K2B	+ 1082.50	1787.15	1.3e-91	2178	D136475 Drosophila melanog
gb.inl:DM08K2B	+ 1080.50	1787.15	4.4e-91	2806	D03646 Fruitfly mRNA for DT
gb.inl:DM08K2B	+ 1067.50	1776.40	2.8e-90	2410	D11441 Drosophila melanogas
gb.inl:DM011441	+ 1066.00	1761.71	3.5e-90	2413	D133790 Drosophila melanog
gb.inl:DM08K4A	+ 1063.00	1756.91	6.4e-90	2361	L133476 Drosophila melanog
gb.inl:DM005209	+ 1061.00	1753.26	1.0e-89	2425	AF03209 Drosophila melanog
gb.inl:AF031357	+ 1061.00	1753.26	2.4e-89	1896	AF031357 Schistosoma manso
gb.inl:AF031358	+ 983.00	1620.86	9.8e-65	466	U63318 Rattus norvegicus hab
gb.ro:NM053318	+ 763.00	1304.71	9.8e-65	750	U60420 Ratius norvegicus acti
gb.prl:PH060420	+ 755.00	1252.56	7.9e-62	2441	AF031391 Brugia pahangi TGF
gb.inl:AF0695300	+ 704.00	1154.19	2.4e-56	3664	AF0695300 Danio rerio activ1
gb.ov:AF0695300	+ 686.50	1119.56	2.0e-54	2615	AF001406 Carassius auratus
gb.ov:AF0695300	+ 679.00	1111.49	5.7e-54	2615	X11223 Gallus gallus activ1
gb.ov:G5U014263	+ 679.00	1093.97	5.4e-53	1903	S70930 actin receptor [Xen
gb.ov:G5U014263	+ 660.50	1081.77	3.5e-52	2315	U57037 Bos taurus actin r
gb.ov:570930	+ 657.50	1079.27	3.5e-52	1833	U57037 Bos taurus actin r
gb.ov:570930	+ 655.00	1071.53	9.6e-52	2542	M88594 Xenopus laevis activ
gb.com:BU057707	+ 655.00	1070.77	1.1e-51	1587	X77533 H.sapiens mRNA for a
gb.ov:KMLAKR	+ 650.50	1068.29	1	1708	M84120 mouse actin r
gb.prl:HSACTIIRE	+ 648.00	1061.92	3.3e-51	2075	S49438 actin receptor [Xen
gb.ro:NM05ACTR2B	+ 648.00	1061.92	3.3e-51	2146	AF104017 Caenorhabditis acti
gb.ov:SA49438	+ 644.50	1059.34	4.6e-51	1542	D31889 Chicken mRNA for ac
gb.inl:DM040117	+ 644.50	1057.44	5.8e-51	1573	X63128 H.sapiens mRNA for a
gb.ov:CHAKATITIA	+ 642.50	1055.31	7.7e-51	1640	L21717 Bos taurus (clones F
gb.prl:HSACTREC					
gb.com:BOVACTRIT					

seq_documentation_block:

Unclassified.
1 (bases 1 to 1482)
Thomson, C. F., Ryden, M., and Jornvall, H.

AUTHORS LAUREZ/C.O. / 21/00/00

